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March 29, 2002

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**BOX PCT**Commissioner for Patents  
Washington, D.C. 20231PCT/JP00/06804  
-filed September 29, 2000

Re: Application of Kei YAMANA, Yukimi NAGASAWA, Hitoshi WADA and Yoshinori KASAHARA  
A NOVEL POLYPEPTIDE AND GENE ENCODING THE SAME  
Assignee: **TELJIN LIMITED**  
Our Ref: Q69170

Dear Sir:

The following documents and fees are submitted herewith in connection with the above application for the purpose of entering the National stage under 35 U.S.C. § 371 and in accordance with Chapter II of the Patent Cooperation Treaty:

- ☒ an executed Declaration and Power of Attorney.
- ☒ an English translation of the International Application.
- ☒ ten (10) sheets of drawings.
- ☒ an executed Assignment and PTO 1595 form.
- ☒ twenty (20) pages of Sequence listing and a 3.5" disk containing Sequence Listing.
- ☒ Statement to Support Filing and Submission in Accordance with 37 C.F.R. §§ 1.821-1.825.
- ☒ International Search Report, Information Disclosure Statement a Form PTO-1449.

It is assumed that copies of the International Application, the International Search Report, the International Preliminary Examination Report, and any Articles 19 and 34 amendments as required by § 371(c) will be supplied directly by the International Bureau, but if further copies are needed, the undersigned can easily provide them upon request.

Applicants note that the second inventor's name changed from Yukimi Takahashi to Yukimi Nagasawa due to marriage. A Declaration is being prepared in this regard and will be filed with a Petition after it is executed.

The Government filing fee is calculated as follows:

Total claims	69	-	20	=	49	x	\$18.00	=	\$882.00
Independent claims	6	-	3	=	3	x	\$84.00	=	\$252.00
Base Fee									\$890.00
Multiple Dependent Claim Fee									\$280.00
<b>TOTAL FILING FEE</b>									<b>\$2304.00</b>
<b>Recordation of Assignment</b>									<b>\$ 40.00</b>
<b>TOTAL FEE</b>									<b>\$2344.00</b>

Checks for the statutory filing fee of \$2304.00 and Assignment recordation fee of \$40.00 are attached. You are also directed and authorized to charge or credit any difference or overpayment to Deposit Account No. 19-4880. The Commissioner is hereby authorized to charge any fees under 37 C.F.R. §§ 1.16, 1.17 and 1.492 which may be required during the entire pendency of the application to Deposit Account No. 19-4880. A duplicate copy of this transmittal letter is attached.

Priority is claimed from:

<u>Country</u>	<u>Application No</u>	<u>Filing Date</u>
Japan	11-275947	September 29, 1999

Respectfully submitted,

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SJM/amt

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 IC10 Rec'd PCT/PTO 29 MAR 2002  
 Washington, DC 20037-3213  
 T 202.293.7060  
 F 202.293.7860

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TN-H822

IC10 Rec'd PCT/PTO 29 MAR 2002

## Description

### A Novel polypeptide and gene encoding the same

#### 5 Field of the Invention

The present invention relates to a novel human, mouse and rat polypeptide having a homology in the amino acid sequence with chondromodulin-I (ChM-I) that is known to have an effect of controlling the growth and  
10 differentiation of chondrocytes and inhibiting angiogenesis, and a human, mouse and rat gene (hereinafter referred to as "ChM1L gene") encoding the same.

#### 15 Background Art

Almost all the bones of mammals are formed through a mechanism called "endochondral bone formation" in which chondrocytes calcify via the growth and differentiation thereof, and are finally replaced with bone. It is known  
20 that a variety of hormones and growth factors participate in this series of process, including insulin-like growth factor (IGF1, IGF2), fibroblast growth factor (FGF), transforming growth factor (TGF), growth hormone and the like. Hiraki et al. isolated ChM-I gene as a factor, in  
25 addition to the above hormones and growth factors, that facilitates the growth and differentiation of chondrocytes (Biochem. Biophys. Res. Commun., 175, 971-977, 1991, European Patent Publication No. 473080). Human ChM-I is synthesized as a type II membrane protein  
30 comprising 334 amino acid residues and, after glycosylation, undergoes processing with a result that the C-terminal portion comprising 120 amino acid residues are extracellularly secreted (Hiraki et al., Eur. J. Biochem. 260, 869-878, 1999). ChM-I not only promotes  
35 the growth of cultured chondrocytes but potently promotes proteoglycan synthesis and the colony formation of chondrocytes in agarose (Inoue et al., Biochem. Biophys.

- 2 -

Res. Commun., 241, 395-400, 1997). ChM-I also promotes the growth of osteoblasts (Mori et al., FEBS Letters, 406, 310-314, 1997).

On the other hand, it has long been pointed out that cartilage remains not only avascular but resistant to infiltration of blood vessels. Hiraki et al. attempted to purify a growth inhibiting factor for vascular endothelial cells from the extracts of cartilaginous tissue, and have succeeded in the complete purification thereof. As a result, it was found to be ChM-I (Hiraki et al., FEBS Letters, 415, 321-324, 1997; Hiraki et al., J. Biol. Chem., 272, 32419-32426, 1997). Generally, the cartilaginous tissue is characterized by being avascular, but in the replacement to the bone tissue, it is believed, infiltration of blood vessels into the cartilaginous tissue is required. In the scheduled region of vascular invasion, the hypertrophy of cartilaginous tissue and the calcification of cartilage matrix occur prior to vascular invasion to be ready for forming the primary point of ossification. In the region where the hypertrophic cartilage and the subsequent ossified cartilage appear, the expression of ChM-I dramatically decreases. Thus, although the expression of the ChM-I gene is cartilage-specific, it is limited to the avascular cartilage that is resistant to vascular invasion. As described above, it is believed that ChM-I not only promotes the growth, differentiation, and maturing of cartilage but inhibits the infiltration of blood vessels by inhibiting the growth of vascular endothelial cells. Thus, the expression in the avascular cartilage and the disappearance of expression in the ossified layer prior to vascular invasion are in good agreement with the bifunctional effect of ChM-I.

In the cartilaginous tissue, bFGF that is a potent angiogenic factor is accumulated in pericellular space in large quantities, and it has been elucidated that ChM-I is present in interterritorial space in such a way as to

surround bFGF (Hiraki et al., J. Biol. Chem., 272, 32419-32426, 1997). Thus, in the avascular cartilage, ChM-I is present in a form that masks the angiogenic factor, and it is thought that the angiogenenic effect of ChM-I may account for the absence of blood vessels in the cartilage (Tanpakusitsu kagaku koso, Vol. 40, No. 5, 1995). It has also been confirmed that ChM-I suppresses the growth of tumor cells by inhibiting the infiltration of blood vessels into human tumor cells in vivo (Hayami et al., FEBS Letters, 458, 436-440, 1999). The expression analysis of ChM-I in various mouse tissues revealed that ChM-I is expressed in the eye and the thymus in addition to the cartilage, but the function of ChM-I in these tissues has yet to be elucidated (Shukunami et al., Int. J. Dev. Biol. 43, 39-49, 1999).

The growth and the expression of differentiation function of chondrocytes plays an important role in the healing process from fracture or various cartilage diseases. Thus ChM-I, a factor that promotes the growth and differentiation of chondrocytes, is a promising candidate for an agent that promotes the growth of chondrocytes (Kokai (Japanese Unexamined Patent Publication) No. 7-138295). In the growth or metastasis of tumor cells, infiltration of blood vessels into tissues is required to obtain energy necessary therefor. Therefore ChM-I that has an effect of inhibiting angiogenesis is also a likely candidate for an anti-cancer agent (Kokai (Japanese Unexamined Patent Publication) No. 7-138295). As described above, ChM-I not only controls the growth and differentiation of chondrocytes but inhibits angiogenesis, and hence its application into drugs is being awaited.

In recent years, biotechnology has made rapid progress, and in association with the development of the human genome project as well, a great number of new genes are being cloned. It is said that the number of human genes amounts to about 100,000, and among the genes



the functions of each of these molecules were analyzed and the similarity and the difference between them were elucidated.

Molecules of the TNF family have the structure of type II membrane proteins and since many of them are expressed mostly in the blood system and the lymphatic system, they have a lot in common in terms of experimental techniques and samples. It is therefore expected that when a new gene belonging to the TNF family was discovered, the speed at which its function was analyzed must have been faster than the molecules discovered earlier. Thus, the discovery of a novel gene having a homology in the amino acid sequence and the analysis of its function would not only facilitates the functional analysis of novel genes to be discovered in the future but the result of analysis permits its comparison with the existing molecules, and therefore it is expected that more detailed findings on the functions of the existing molecules could be obtained.

Generally, when a novel gene encoding a protein having a homology in the amino acid sequence with existing molecules is cloned, the techniques and materials to be used for functional analysis may be referred to the examples of the existing molecules. However, even a molecule having a homology in the amino acid sequence is thought to have its own unique function as in the above-mentioned TNF family, and thus when its application into pharmaceutical products is envisaged, it is necessary to demonstrate the expression and purification of the recombinant protein, the generation of antibody, the expression of mRNA and protein at various tissues and the like, and thereby to clarify the difference in the structure and function from the existing molecules.

#### Disclosure of the Invention

Thus, it is an object of the present invention to

- 6 -

provide a new polypeptide similar to ChM-I and a gene encoding the same. It is also an object of the present invention to implement the generation of antibody against said polypeptide, the analysis of expression levels of  
5 said gene and the polypeptide, the expression and structural analysis of the recombinant protein and the like in order to clarify its similarity and difference with ChM-I, and to elucidate the function so as to enable the elucidation of pathological states, diagnosis,  
10 treatment etc. of diseases in which they are involved.

ChM-I is a type II membrane protein that regulates the growth and differentiation of chondrocytes and inhibits angiogenesis, and is a promising candidate for application into pharmaceutical products. Thus, once a  
15 gene encoding a new polypeptide similar to ChM-I has been provided, it is believed, its expression level in various cells and its structure and function can be analyzed, and the analysis of the expression products would enable the elucidation of pathology, diagnosis and treatment etc. of  
20 diseases in which they are involved. At present, however, there are no reports on molecules having a homology with the amino acid sequence of ChM-I, and it is unknown whether ChM-I forms a gene family or not. Thus, if a new polypeptide similar to ChM-I and a gene encoding  
25 the same are shown to be present, the analysis of structure, function etc. thereof would permit the study on its similarity and difference with ChM-I, which in turn would accelerates the elucidation of physiological functions of the molecules with one another, the  
30 elucidation of pathological states in which these molecules are involved, diagnosis, the development of therapeutic agents and the like.

After intensive and extensive research to attain the above purposes, the inventors of the present invention  
35 have succeeded in isolating a gene (ChM1L gene) that meets the above purposes, from human, mouse and rat cDNA libraries, and carried out the analysis of its expression

- 7 -

level in various tissues, the generation of an antibody against said polypeptide, the expression of a polypeptide encoded by said gene in a mammalian cell, its detection and purification and the like, demonstrating that said  
5 polypeptide has an effect of inhibiting angiogenesis, and we hereby have completed the present invention.

Thus, the present invention is a gene encoding a polypeptide that substantially comprises the amino acid sequence as set forth in SEQ ID NO: 2, 4, and 6. As the  
10 above gene, there can be mentioned the nucleotide sequence represented by SEQ ID NO: 1, 3, and 5.

Furthermore, the present invention is a polypeptide encoded by a human, mouse and rat gene which polypeptide substantially comprises the amino acid sequence as set  
15 forth in SEQ ID NO: 2, 4 and 6.

The present invention is also an oligonucleotide probe that hybridizes to at least part of the above gene.

The present invention is also a recombinant DNA comprising the above gene.

20 The present invention is also a transformant transformed with the above recombinant DNA.

The present invention is also a method of producing the above polypeptide which method comprises culturing the above transformant and harvesting a polypeptide  
25 encoded by the gene of the present invention from the culture.

The present invention is also a monoclonal antibody or polyclonal antibody that specifically reacts with the above polypeptide.

30 The present invention is also a hybridoma that produces the above monoclonal antibody that is obtained by fusing an antibody-producing cell immunized with the above polypeptide to a myeloma cell.

The present invention is also a reagent for  
35 detecting genes said reagent comprising the above oligonucleotide probe.

The present invention is also a diagnostic kit that



- 8 -

comprises the above polypeptide and the above monoclonal antibody or polyclonal antibody.

5 The present invention is also a pharmaceutical composition comprising a polypeptide encoded by the gene that substantially comprises the amino acid sequence as set forth in SEQ ID NO: 2, 4, or 6.

10 The present invention is also a pharmaceutical composition comprising a monoclonal antibody or a polyclonal antibody that specifically reacts with the above polypeptide.

The present invention is also a pharmaceutical composition comprising an antisense oligonucleotide that specifically hybridizes to part of the above gene.

15 The present invention is also a pharmaceutical composition comprising a nucleic acid that can be used in gene therapy said composition comprising at least part of the above gene.

The present invention is also a polypeptide wherein the above polypeptide is a membrane-bound form.

20 The present invention is also a gene encoding the above membrane-bound polypeptide.

The present invention is also a gene wherein the above human gene is present on chromosome X.

25 The present invention is also a polypeptide wherein the above polypeptide has an effect of inhibiting angiogenesis.

The present invention is also a gene encoding the above polypeptide that has the above effect of inhibiting angiogenesis.

30

#### Brief Explanation of the Drawings

Figure 1A is a result in which the homology of amino acid sequences of human ChM1L and human ChM-I were compared.

35 Figure 1B is a result in which the homology of amino acid sequences of human, mouse and rat ChM1L were compared.



- 10 -

Figure 7 shows the result in which the expression of ChM1L protein in mouse rib cartilage was detected by immunostaining using anti-ChM1L polypeptide antibody.

Figure 8 shows the result in which a soluble human  
 5 ChM1L protein that was expressed in the culture liquid of COS7 cells was purified by affinity chromatography using anti-FLAG M2 affinity gel, electrophoresed, and stained with Coomassie brilliant blue. Lane 1 shows the result of electrophoresis of the culture supernatant of COS7  
 10 cells and lane 2 shows that of the purified ChM1L protein.

Figure 9 shows the result in which the tube-like structure-forming system of the human umbilical vein endothelial cells were treated with (a) the buffer alone,  
 15 (b) 20 µg of bovine serum albumin (BSA), (c) 10 µg of soluble human ChM1L, (d) 20 µg of soluble human ChM1L, (e) 1 µg of platelet factor 4 (PF-4), and (f) 10 µg of PF-4.

## 20 Best Mode for Carrying Out the Invention

In accordance with the present invention, "substantially comprise" means that the gene or polypeptide of the present invention, as long as it retains its function, may have mutations such as  
 25 substitution, insertion, or deletion in the nucleotide sequence as set forth in SEQ ID NO: 1, 3, or 5 or the amino acid sequence as set forth in SEQ ID NO: 2, 4, or 6.

The ChM1L gene sequence of the present invention may  
 30 be obtained by the RACE method (RACE: Rapid amplification of cDNA ends; Frohman, M. A. et al., Proc. Natl. Acad. Sci. USA, 85, 8998-9002, 1988), the outline of which method is as follows:

Generally, the RACE method enables one to obtain a  
 35 full-length cDNA in an efficient manner, when a portion of a cDNA sequence is known. Primers are constructed

- 11 -

from known sequence regions to permit elongation in each of the 3'-end or 5'-end direction, and then cDNA is amplified by the polymerase chain reaction (PCR, Science, 230, 1350-1354, 1985). When a PCR method is carried out, 5 primers that specifically anneal are used in the known region, and primers that anneal to the sequence tagged by a ligation reaction are used in the 3'-end and the 5'-end. Thus, the regions amplified by PCR contain unknown regions. The isolation and purification of the amplified 10 cDNA fragment can be performed according to a standard method as described below, for example gel electrophoresis may be employed. The determination of nucleotide sequence of the DNA fragment thus obtained may be performed according to a standard method such as the 15 dideoxy method (Proc. Natl. Acad. Sci. USA, 74, 5463-5467, 1977) and the Maxam-Gilbert method (Methods in Enzymology, 65, 499, 1980). Such determination of nucleotide sequences may also be carried out using commercially available sequencing kits, etc.

20 More specifically, it is outlined as follows, though more detailed explanations thereof will be made hereinafter in Example 2. Using the amino acid sequence of human ChM-I, TBLASTN search was performed for the EST data base (dbEST, EST: Expressed sequence tag) in the DNA 25 data bank of Japan (DDBJ) to detect an EST file, Genbank accession number AI123839. AI123839, which is a nucleotide sequence fragment registered in dbEST, was found for the first time by the above TBLASTN search to be a novel gene fragment encoding an amino acid sequence 30 similar to ChM-I. Thus, primers were synthesized from part of the sequence of cDNA obtained from dbEST, and the sequence of human ChM1L gene was determined using the RACE method. Subsequently, the sequences of mouse and rat ChM1L genes were similarly determined. The sequences 35 of human, mouse and rat ChM1L genes are shown in SEQ ID NO: 1, 3 and 5, and the amino acid sequences of the peptides encoded thereby are shown in SEQ ID NO: 2, 4 and

6.

The polypeptides encoded by the ChM1L genes of the present invention are composed of 317 amino acids (SEQ ID NO: 2, 4 and 6). The amino acid sequence of ChM1L has a  
5 homology with ChM-I, in particular a very high homology with the C-terminal portion that is extracellularly secreted after the processing of ChM-I (Figure 1(a)). The amino acid sequences of ChM1L have a very high homology in between humans, mice, and rats (Figure 1(b)).  
10 From the hydrophobicity analysis of the amino acid sequence, ChM1L similarly to ChM-I is thought to be a molecule having the structure of the type II membrane proteins (Figure 2). As shown in Figure 2, in both of said polypeptide and ChM-I, a hydrophobic domain  
15 comprising about 20 amino acids that is uniquely found in molecules having a membrane-binding activity is present in the vicinity of several dozen amino acids from the N-terminal. That said polypeptide is a molecule having the type II membrane protein structure was also demonstrated  
20 by the result in Example 8 in which said polypeptide was expressed in COS7 cells (Figure 4).

The human ChM1L gene of the present invention was shown, as described below in Example 12, to be present on chromosome X (Genbank accession No. AL035608).

25 As the ChM1L gene of the present invention, there can be mentioned cDNA, chemically synthesized DNA, DNA isolated by PCR, genomic DNA and combinations thereof. Using a standard method, said genomic DNA may also be isolated by hybridization with the ChM1L gene that is  
30 disclosed herein. RNA that was transcribed from said ChM1L gene is also encompassed in the present invention. The sequences of the gene of the present invention represented by SEQ ID NO: 1, 3 and 5 are a combination example of codons representing amino acid residues  
35 encoded by them. The ChM1L gene of the present invention is not limited to this, and it is also possible to have a DNA sequence obtained by combining codons to amino acid

- 13 -

residues and then selected. The selection of said codon may be performed according to a standard method, and for example the frequency of use of host's codons to be used is taken into account (Nucleic Acids Research, 9, 43-74, 1981).

Furthermore, the ChM1L gene of the present invention encompasses DNA sequences encoding mutants in which parts of the amino acid sequence as set forth in SEQ ID NO: 2, 4, and 6 are substituted, deleted, or added. The production, modification (mutation) and the like of these polypeptides may naturally occur, and can be obtained by post-translational modification or by a genetic engineering method such as site-specific mutagenesis (Methods in Enzymology, 154, 350, 367-382, 1987; *ibid.*, 100, 468, 1983; Nucleic Acids Research, 12, 9441, 1984; Zoku Seikagaku Jikken Koza 1 (Sequel to Biochemistry Experimental Series 1) "Idensi Kenkyuuhou II (Gene Study Method II)", edited by The Japanese Biochemical Society, 105, 1986) and the like.

The production of the ChM1L gene of the present invention may be readily performed by a common genetic engineering method based on the sequence information of the ChM1L gene of the present invention (Molecular Cloning, 2nd Ed., Cold Spring Harbor Laboratory Press, 1989; Zoku Seikagaku Jikken Koza (Sequel to Biochemistry Experimental Series) "Idensi Kenkyuuhou I, II, III (Gene Study Method I, II, III)", edited by The Japanese Biochemical Society, 1986).

This can be attained by, for example, selecting the desired clones from a cDNA library (prepared from a suitable cell source that expresses the ChM1L gene according to a standard method) using suitable probes and antibodies unique to the gene of the present invention (Proc. Natl. Acad. Sci. USA, 78, 6613, 1981; Science, 222, 778, 1983, and the like).

In the above method, as the cell source, there can be illustrated various cells, tissues, and cultured cells









- 17 -

reduce the expression of the ChM1L gene, and the like are considered to be used as therapeutic agents for these diseases. The agonists and antagonists mentioned above are intended to include peptides, proteins, low molecular weight compounds and the like, but the physical properties are not limited to them as long as they retain the function.

In various tissues of a fetus, the expression of ChM1L mRNA was noted in the eyeball, the kidney, the stomach, the whole rib and the trachea (Figure 3(b)). In an adult mouse, ChM1L mRNA was not expressed in the kidney or the stomach, but the ChM1L mRNA was expressed in these tissues of the fetus suggests that ChM1L is involved in the development and morphogenesis of these organs. Thus, ChM1L is thought to be also associated with the repair and regeneration of these organs in adults. It was also revealed that ChM1L mRNA is expressed in the trachea. Thus, the ChM1L gene and the ChM1L polypeptide of the present invention, antagonists and agonists to ChM1L including antibody that binds to ChM1L, agents that promote or reduce the expression of the ChM1L gene, and the like are considered to be used as therapeutic agents for kidney-related diseases such as chronic kidney failure, stomach-related diseases such as gastric cancer and gastric ulcer, and trachea-related respiratory diseases such as chronic bronchitis and asthma.

In the developmental stage of the fetus, the expression of ChM1L mRNA is very weak at day 10 of gestation, and the expression increases at around days 11 to 13 (Figure 3(c)). On the other hand, though ChM-I, similarly to ChM1L, increases in expression with the development of the fetus, it was evidently expressed more strongly than ChM1L on days 10 and 11 of gestation. It is, therefore, clear that ChM1L lags behind ChM-I in the expression, and that these molecules have different functions in the development of the fetus. The increase









- 22 -

detect ChM1L recombinant proteins. No expression of ChM1L recombinant proteins was confirmed in the culture supernatant, whereas in the cellular components two bands were detected at around 40 kDa.

5           Thus, it was revealed that the ChM1L protein is a membrane-bound protein. On the other hand, it has been confirmed that when ChM-I is expressed in COS7 cells, it is secreted as a soluble protein in the culture supernatant (Hiraki et al., J. Biol. Chem., 272, 32419-  
10           32426, 1997). Thus the analysis with COS7 cells revealed that ChM1L and ChM-I are proteins having different structures. That is, it was shown that ChM1L is a cell membrane-bound protein, while ChM-I is a secretory protein, and that the processing mechanisms of these  
15           molecules are different. Among the two bands for the ChM1L protein, the band at the high molecular weight was found to be a form modified by a N-linked sugar chain in the Example 10 described below (Figure 6).

20           The ChM1L protein thus expressed can affinity purified using a ChM1L-specific antibody or an antibody against the tag (His tag) in which 6 residues of histidine are fused, a nickel column and the like.

25           The polypeptide encoded by the ChM1L gene of the present invention may be any of a membrane-bound polypeptide and a soluble polypeptide having no property of binding to the cell membrane. For example, there may be cases in which after the polypeptide is expressed as a membrane-bound polypeptide on the cell membrane, it is cleaved to become a soluble polypeptide. Though the  
30           ChM1L protein was detected as a membrane-bound protein in the expression in COS7 cells (Example 8), it may undergo processing thereby to be a soluble protein when the host cell or the culture condition is different. Furthermore, the soluble polypeptide that lacks the transmembrane  
35           domain can be expressed by fusing a heterologous signal peptide to the N-terminal.

More specifically, the method of expressing the

- 23 -

soluble ChM1L protein is outlined as follows, though more detailed explanations thereof will be made hereinafter in Example 9.

5 A vector was constructed that has integrated, into pCAGGS vector, a nucleotide sequence encoding a protein in which the signal sequence of preprotrypsin, a FLAG tag, the C-terminal end of the extracellular region of ChM1L were fused from the N-terminal end (Example 5).  
10 The ChM1L protein that was expressed using this vector was secreted into the culture liquid as a soluble protein after the signal sequence of preprotrypsin was cleaved (Example 9, Figure 5).

The soluble ChM1L polypeptide thus secreted into the culture liquid can be purified using anti-ChM1L antibody  
15 or anti-FLAG antibody (Sigma) because a FLAG tag is fused thereto. It is also possible to remove the FLAG tag by cleaving the FLAG fusion protein with enterokinase.

More specifically, the method of purifying the soluble ChM1L protein is outlined as follows though more  
20 detailed explanations thereof will be made hereinafter in Example 13.

Using a Lipofectamine reagent (GIBCO BRL) according to the instruction attached to the product, pSF-shChM1L was transfected into COS7 cells, and 48 hours later the  
25 culture supernatant was harvested. From this culture supernatant, a soluble ChM1L protein was purified by affinity chromatography using anti-FLAG M2 affinity gel (Sigma) (Figure 8).

The ChM1L polypeptide of the present invention can  
30 be used as a polypeptide-purifying reagent. Said polypeptide bound to a solid support is very useful for the purification of polypeptides that can bind to said peptide by affinity chromatography. As the polypeptide that can bind to the ChM1L polypeptide, there may be  
35 illustrated soluble polypeptides, membrane-bound polypeptides, antibodies and the like. The soluble ChM1L polypeptide may be readily used for the addition into the



cell culture liquid in vitro, or intravenous administration in vivo.

Using the polypeptide encoded by the ChM1L gene of the present invention, a specific antibody can be generated. An antigen as used herein includes a polypeptide produced in large quantities according to the above genetic engineering method or a chemically synthesized polypeptide, and an antibody obtained may be any of polyclonal antibody or monoclonal antibody, and can be effectively used for the purification, measurement, recognition, and the like of said polypeptide. Hence, polyclonal antibodies and monoclonal antibodies against said polypeptide can be used for treatment or the development of therapeutic methods for diseases that are mediated (directly or indirectly) by said polypeptide, and can also be used as diagnostic reagents for the above diseases.



- 26 -

or lymphatic vessels, the perichondrium is thought to regulate the infiltration of blood vessels, nerves, or lymphatic vessels into the chondrocytes since the perichondrium is present at the interface of  
 5 cartilaginous tissues and other tissues. Thus, though perichondrium is recognized to be an important tissue, it has no definite definition and detailed study has not been made at present. One reason for this that there are no molecules that are perichondrium-specifically  
 10 expressed have been elucidated at all.

Accordingly, if the presence of molecules that are specifically expressed in the tissue termed as perichondrium surrounding the cartilaginous tissue is demonstrated, it would provide a very important tool in  
 15 the study of perichondrium and cartilaginous tissues.

The ChM1L of the present invention is the only molecule that was demonstrated to be perichondrium-specifically expressed, and is thought to regulate the infiltration of blood vessels, nerves, or lymphatic  
 20 vessels into the cartilaginous tissues.

Hence, the discovery of the ChM1L gene and the result of functional analysis of ChM1L included herein is believed to provide, from now on, a new perspective on the etiology and the development of therapeutic methods  
 25 for diseases in which perichondrium and other ChM1L-expressing tissues including the cartilaginous tissue are involved.

Thus, the ChM1L gene and the ChM1L polypeptide of the present invention, antagonists and agonists to ChM1L including antibody that binds to ChM1L, and agents that  
 30 promote or reduce the expression of the ChM1L gene are considered to be used as therapeutic agents for diseases in which the above ChM1L-expressing cells are involved.

The above expression analysis of mRNA and the result of immunostaining revealed that the ChM1L gene and the polypeptide encoded thereby of the present invention are  
 35 expressed in the brain, the eyeball, the skeletal muscle,

- 27 -

the thyroid, the whole rib including cartilage, the kidney, the stomach, the trachea, and cells that assume a fibroblast-like flat form occurring in such a way as to surround the cartilaginous tissue. It suggests, therefore, that the ChM1L gene and the polypeptide encoded thereby of the present invention may be involved in diseases associated with the above tissues that have been confirmed to express them, such as diabetic retinopathy, muscular dystrophy, Basedow's disease, chronic kidney failure, stomach cancer, chronic bronchitis, osteoarthritis and rheumatoid arthritis.

Hence, the ChM1L gene and the ChM1L polypeptide of the present invention, antagonists and agonists to ChM1L including antibody that binds to ChM1L, and agents that promote or reduce the expression of the ChM1L gene are considered to be used as therapeutic agents for these diseases.

#### Examples

The present invention will now be explained more specifically with reference to the following examples. It should be noted, however, that the present invention is not limited to these examples.

#### Example 1. Analysis of the ChM1L amino acid sequence

The homology of amino acid residues of ChM-I and ChM1L was compared (Figure 1(a)). The amino acid sequence was represented by one alphabetical letter. ChM1L has a homology with ChM-I throughout the molecule, but it was found that ChM1L has a particularly high homology with the C-terminal of ChM-I that is extracellularly secreted following the processing of ChM-I.

The homology of amino acid sequences of human, mouse and rat ChM1L was compared (Figure 1(b)). The ChM1L polypeptide is composed of 317 amino acids in humans, mice, and rats, but the 300 amino acid residues were identical in the three (about 95%).



determined using a DNA sequencer (ABI PRISM (TM) 310 Genetic Analyzer) of PE Applied Biosystems and ABI PRISM (TM) BigDye Terminator Cycle Sequencing Ready Reaction kit.

5           The nucleotide sequence of human ChM1L cDNA is shown in SEQ ID NO: 1 and its amino acid sequence is shown in SEQ ID NO: 2.

          Since the amino acid sequence encoded by the human ChM1L gene represented by SEQ ID NO: 1 has a homology  
10       with human ChM-I, the gene was decided to be termed as the ChM1L gene (ChM-I like gene).

          The coding sequence (CDS) of Human ChM1L cDNA was amplified by PCR, electrophoresed on agarose, and then was purified, which was then cloned using pCR-Script (TM)  
15       Amp cloning kit (Stratagene) according to the instruction attached to the product. The sequence of the primers used in PCR are shown in SEQ ID NO: 7 (forward primer) and SEQ ID NO: 8 (reverse primer). The ChM1L gene sequence that has been integrated into the vector was  
20       determined using ABI PRISM (TM) 310 Genetic Analyzer of PE Applied Biosystems and ABI PRISM (TM) BigDye Terminator Cycle Sequencing Ready Reaction kit.

          Using the amino acid sequence (SEQ ID NO: 2) of human ChM1L, TBLASTN search was carried out as described  
25       in the above human case. As a result, as a gene fragment that encodes mouse ChM1L, EST file, Genbank accession number AV009191 was detected, and as a gene fragment that encodes rat ChM1L, EST file, Genbank accession number AI112003 was detected. Using Mouse 11-day Embryo  
30       Marathon-Ready (TM) cDNA and Rat Skeletal muscle Marathon-Ready (TM) cDNA by Clontech, sequences of mouse and rat ChM1L genes were determined by the RACE method as described in the isolation of the human ChM1L gene.

          The nucleotide sequence of mouse ChM1L cDNA is shown  
35       in SEQ ID NO: 3 and the amino acid sequence is shown in SEQ ID NO: 4. The nucleotide sequence of rat ChM1L cDNA is shown in SEQ ID NO: 5 and the amino acid sequence is







- 32 -

pCAGGS vectors containing the hChM1L, mChM1L, hChM1LHis and mChM1LHis: pCAGGS-hChM1L, pCAGGS-mChM1L, pCAGGS-hChM1LHis, and pCAGGS-mChM1LHis

Example 5. Construction of vectors that express human soluble ChM1L protein to which a FLAG tag is fused

The FLAG tag (Sigma) as used herein is a hydrophilic marker peptide comprising eight amino acids (Asp Tyr Lys Asp Asp Asp Asp Lys), and the last five amino acids (Asp Asp Asp Asp Lys) is a recognition sequence for enterokinase. The vectors constructed in this Example can express a protein in which the signal sequence of preprotrypsin, a FLAG tag, the C-terminal end of the extracellular region of ChM1L were fused from the N-terminal end. The protein that was expressed using this vector is secreted into the culture liquid as a soluble protein after the signal sequence of preprotrypsin is cleaved, as explained in detail hereinafter in Example 9. Since a FLAG tag is fused to the protein expressed with this vector, the protein can be purified using anti-FLAG antibody (Sigma) and by cleaving the fusion protein with enterokinase, the FLAG tag can also be removed.

A vector was constructed in which a nucleotide sequence (SEQ ID NO: 19, contained in pFLAG-CMV-1 vector manufactured by Sigma) encoding the signal sequence of preprotrypsin and a FLAG tag (SEQ ID NO: 20) from the N-terminal was integrated into the pCAGGS vector (hereinafter referred to as pSF vector). A nucleotide sequence (nucleotide sequence No. 684 to 1020 of SEQ ID NO: 1) encoding the amino acids No. 212 to 317 of human ChM1L represented by SEQ ID NO: 2 and the translation termination codon was amplified by the PCR method, and the amplified product was integrated into the 3'-end of the nucleotide sequence encoding the FLAG tag of the pSF vector. The sequences of primers used for PCR are shown in SEQ ID NO: 21 (forward primer) and SEQ ID NO: 8 (reverse primer). The integration of the sequence of interest into the constructed vector was confirmed using

- 33 -

a ABI PRISM (TM) 310 Genetic Analyzer (PE Applied Biosystems) and ABI PRISM (TM) BigDye Terminator Cycle Sequencing Ready Reaction kit according to the instruction attached to the product. The nucleotide  
 5 sequence of this Example integrated into the vector is shown in SEQ ID NO: 22, and the amino acid sequence encoded thereby is shown in SEQ ID NO: 23. The vector constructed in this Example will be abbreviated to pSF-shChM1L.

10 Example 6. Expression analysis of ChM1L mRNA  
Expression analysis of ChM1L mRNA in various tissues of an adult (10-week old): Figure 3(a)

A 10-week old C57BL/6 mouse was dissected and each tissue was extracted, which was immediately frozen in  
 15 liquid nitrogen. The frozen tissue was ground into small pieces, and using IOSGEN (Nippon Gene) according to the instruction attached to the product the total RNA of each tissue was obtained. With one µg of total RNA of each tissue as template, 20 µl of cDNA was synthesized using  
 20 Superscript II preamplification kit (GIBCO BRL) according to the instruction attached to the product. In RT-PCR, the total liquid volume of the reaction system was set at 50 µl, and for each tissue 0.5 µl of cDNA, 0.25 µl of ExTaq polymerase (Takara Shuzo) were used, to which the  
 25 forward primer (SEQ ID NO: 9) and the reverse primer (SEQ ID NO: 10) were added to a concentration of 0.2 µm.

Using GeneAmp (TM) PCR System 9700 (PE Applied Biosystems), PCR amplification was performed for 30 cycles with each cycle comprising 96°C for 30 seconds,  
 30 60°C for 30 seconds, and 72°C for one minute. The reaction mixture obtained was subjected to electrophoresis on a 1% agarose gel containing ethidium bromide and then the gel was observed under UV irradiation to examine the expression of ChM1L mRNA in  
 35 each tissue.

As shown in Figure 3(a), the expression of ChM1L

mRNA in each tissue of an adult mouse was observed in the brain, the eyeball, the skeletal muscle, the whole rib, and the thyroid. The expression of ChM-I in mice has been confirmed in the eyeball, the thymus, the cartilage, and the whole rib. It is therefore clear that ChM1L and ChM-I are expressed in different tissues in the living body, suggesting that their physiological functions are different.

Expression analysis of ChM1L mRNA in various tissues of a fetus (day 17 of gestation): Figure 3(b)

A fetus of C57BL/6 mouse on day 17 of gestation was removed by Caesarean section. Each tissue was taken out, and was immediately frozen in liquid nitrogen. The extraction of total RNA from the frozen tissue, cDNA synthesis, and RT-PCR were performed as described in the above <Expression analysis of ChM1L mRNA in various tissues of an adult mouse>.

As shown in Figure 3(b), the expression of ChM1L mRNA in each tissue of a fetus mouse was observed in the eyeball, the kidney, the stomach, the whole rib, and the trachea. In the fetus, expression in the kidney and the stomach was observed, in which no expression was observed in the adult mouse. It is therefore likely that ChM1L is involved in the development and morphogenesis of these organs, and is also considered to be involved in the repair and regeneration of organs. It was also revealed that ChM1L mRNA is expressed in the trachea.

Expression analysis of ChM1L mRNA during the developmental stage of a fetus: Figure 3(c)

A fetus of C57BL/6 mouse on each day from day 10 of gestation to childbirth was removed by Caesarean section. Each the whole fetuses was frozen in liquid nitrogen. The extraction of total RNA from the frozen fetus, cDNA synthesis, and the implementation of RT-PCR were performed as described in the above <Expression analysis of ChM1L mRNA in various tissues of an adult mouse>.

The analysis of ChM-I mRNA was carried out using a

- 35 -

fusion protein (SEQ ID NO: 23) and a reverse primer (SEQ ID NO: 24) under the same condition as above.

As shown in Figure 3(c), the expression of ChM1L mRNA during the developmental stage of the fetus is very weak on day 10 of gestation and is increased in expression on days 11 to 13. On the other hand, though the expression of ChM-I was also increased as for ChM1L, it exhibited an evidently stronger expression than ChM1L on days 10 and 11 of gestation. It is therefore clear that the expression of ChM1L lags behind ChM-I during the developmental stage of the fetus, and that these molecules have different functions in the fetus development.

#### Example 7. Generation of anti-ChM1L peptide polyclonal antibody

A peptide having cysteine at the C-terminal of the sequence from 245 to 252 residues shown in SEQ ID NO: 2 of human ChM1L was chemically synthesized. To this synthetic peptide, MBS/KLH (m-maleimidobenzoyl-N-hydroxysuccinimide ester/keyhole limpet hemocyanin, Boehringer Mannheim) was coupled. After the complex was dissolved in physiological saline, an equal amount of Freund's complete adjuvant (FCA) was added, which was sonicated to prepare an emulsion. This emulsion was subcutaneously given to a rabbit as the initial immunization. Four weeks after the initial immunization, a booster immunization was carried out using Freund's incomplete adjuvant (FIA) to the femoral muscle, and thereafter immunization by subcutaneous administration was carried out for four times at an interval of about two weeks or four weeks. During the booster immunization, blood was partially taken from the auricle, and after the final immunization the entire blood was taken and serum was separated. By affinity purification using a peptide column, an anti-ChM1L peptide polyclonal antibody was obtained.

Example 8. Analysis of human and mouse ChM1L recombinant protein by Western blotting: Figure 4

Using the lipofectamine reagent (GIBCO BRL) according to the instruction attached to the product, 5 pCAGGS, pCAGGS-hChM1L and pCAGGS-mChM1L (Figure 4(a) and (c)), or pCAGGS, pCAGGS-hChM1LHis and pCAGGS-mChM1LHis (Figure 4(b) and (d)) were transfected into COS7 cells. About 48 hours after the transfection, the culture supernatant and the cellular components were subjected to 10 sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) on a 12.5% gel, and then transferred to a nitrocellulose membrane. A primary antibody reaction and a secondary antibody reaction were carried out, and then subjected to a color developing reaction using the 15 ECLplus reagent (Amersham Pharmacia Biotech) according to the instruction attached to the product. In the Western blot in which pCAGGS, pCAGGS-hChM1L and pCAGGS-mChM1L were transfected, anti-ChM1L polyclonal antibody described in the above Example was used as the primary 20 antibody and horseradish peroxidase (HRP)-labelled anti-rabbit IgG antibody (DAKO) was used as the secondary antibody, and in the Western blot in which pCAGGS, pCAGGS-hChM1LHis and pCAGGS-mChM1LHis were transfected, anti-His tag antibody (Invitrogen) was used as the 25 primary antibody and HRP-labelled anti-mouse IgG antibody (Amersham Pharmacia Biotech) was used as the secondary antibody.

SDS-PAGE was carried out for the same samples as in Western blot, and the results of staining with Coomassie 30 brilliant blue (CBB) are shown in Figure 4(a) and (b).

As a result of Western blot, no ChM1L band was confirmed in any of the culture supernatants. In cellular components, as shown in Figure 4(b) and (d), recombinant ChM1L protein was detected as two bands at 35 around 40 kDa whether anti-ChM1L peptide antibody or anti-His tag antibody was used. As will be described in detail in the Examples below, the band at the high



ECLplus reagent (Amersham Pharmacia Biotech) was used according to the instruction attached to the product to perform color development reaction.

As shown in Figure 6, the band of ChM1L protein at the high molecular weight disappeared only when treated with pNGase F (lanes 2 and 5). It was therefore demonstrated that the ChM1L protein has been modified with a N-linked sugar chain.

Example 11. Analysis of ChM1L protein at cartilago costalis by immunostaining

An about 10-week old C57BL/6 mouse was dissected to remove the whole rib, which was fixed in a 10 mM phosphate buffer (pH 7.4) (PBS) containing 4% paraformaldehyde, embedded in paraffin, and then sections were prepared. Each step of immunostaining was carried out using Histfine SAB-PO(R) kit (Nichirei) according to the instruction attached to the product, of which outline is as follows: After deparaffinization, endogenous peroxidase was digested with a 3% hydrogen peroxide. After washing with PBS followed by blocking with 10% normal goat serum, the above-mentioned anti-ChM1L peptide antibody at a dilution of 1/160 was added and incubated overnight at 4°C. As a negative control, rabbit IgG was used. After biotin-labelled anti-rabbit IgG antibody and peroxidase-labelled streptoavidin were allowed to react, 3,3-diaminobendizine/4HCl was added to perform a color development reaction. The nucleus was stained with haematoxylin, enclosed, and then observed.

As shown in Figure 7, ChM1L protein is expressed in cells that assume a fibroblast-like flat form occurring in such a way as to surround the cartilaginous tissue. On the other hand, there were no expressions observed in the cartilage cell in which the expression of ChM-I has been reported.

Example 12. Chromosome mapping of the human ChM1L gene

Using the gene sequence (SEQ ID NO: 1) of human ChM1L, BLASTN search was performed for the entire DDBJ

data from the DNA data bank of Japan (DDBJ). As a result, Genbank accession number AL035608 was detected as the genome sequence of the ChM1L gene. AL035608 is a sequence mapped on chromosome X. It is therefore clear that the human ChM1L gene is present on chromosome X.

Example 13. Purification of a soluble human ChM1L recombinant protein

Using a Lipofectamine reagent (GIBCO BRL) according to the instruction attached to the product, pSF-shChM1L was transfected into COS7 cells, and 48 hours later the culture supernatant was harvested. Using anti-FLAG M2 affinity gel (Sigma), an affinity column was prepared, and the culture supernatant was applied to the column. After washing the column three times in 25 mM Tris-HCl, 150 mM NaCl (pH 7.4), it was eluted with 0.1 M glycine-HCl (pH 3.5), and the eluent was neutralized with a 1/20 volume of 1M Tris-HCl (pH 9.5).

The culture supernatant and the eluent were subjected to SDA-PAGE and then were stained with Coomassie brilliant blue (CBB), the result of which is shown in Figure 8. Though there are a variety of proteins in the culture supernatant (Figure 8, lane 1), soluble human ChM1L protein was confirmed as an about 20 kDa band in the eluent. This revealed that soluble human ChM1L protein was concentrated and purified by the above procedure (Figure 8, lane 2).

Example 14. Study on the effect of inhibiting angiogenesis using human umbilical vein endothelial cells

Human umbilical vein endothelial cells (HUVECs, Clonetics) were cultured in a exclusive medium (EGM (TM)-2 Bullet Kit(TM), Clonetics) for endothelial cells. To a 12-well plate, Growth factor reduced Matrigel (Becton Dickinson) was added to 600 µl/well, which was then incubated at 37°C for 30 minutes. Using a heparin-free exclusive medium for endothelial cells diluted 1/8 in the essential medium (EBM (TM)-2, Clonetics) for endothelial cells, a cell suspension containing 5 x 10<sup>4</sup> cells/ml of



- 40 -

HUVECs were prepared.

Each test substance solution was prepared as a solution in which a 1/20 volume of 1M Tris-HCl (pH 9.5) was added to 0.1M glycine-HCl (pH 3.5), and 200  $\mu$ l/well of it was treated. The above buffer and bovine serum albumin (BSA) at 20  $\mu$ g/well as the negative control, platelet factor 4 (PF-4, CHEMICON) at 1 and 10  $\mu$ g/well as the positive control, and the eluted fractions of Example 13 at 10 and 20  $\mu$ g/well as the soluble human ChM1L recombinant protein were treated. Two ml of the cell suspension ( $1 \times 10^5$  cells) and 200  $\mu$ l of the test substance solution were mixed, and seeded into a 12-well plate coated with Growth factor reduced Matrigel. Nine hours later, the formation of tube-like structures was examined and were a photograph was taken. The result is shown in Figure 9. In the negative control, HUVECs formed tube-like structures (Figure 9(a) and (b)), but when ChM1L at 20  $\mu$ g/well (Figure 9(d)) was treated the formation of tube-like structures was inhibited as compared to the negative control.

It was therefore revealed that ChM1L has an effect of inhibiting angiogenesis and thus the soluble ChM1L polypeptide can be used as a therapeutic agent for disease accompanied by angiogenesis such as diabetic retinopathy, cancer, and rheumatoid arthritis.

- 41 -

What is claimed is:

1. A human gene encoding a polypeptide that substantially comprises the amino acid sequence as set forth in SEQ ID NO: 2.
- 5 2. A mouse gene encoding a polypeptide that substantially comprises the amino acid sequence as set forth in SEQ ID NO: 4.
3. A rat gene encoding a polypeptide that substantially comprises the amino acid sequence as set forth in SEQ ID NO: 6.
- 10 4. The human gene according to claim 1 having the nucleotide sequence as set forth in SEQ ID NO: 1.
5. The mouse gene according to claim 2 having the nucleotide sequence as set forth in SEQ ID NO: 3.
- 15 6. The rat gene according to claim 3 having the nucleotide sequence as set forth in SEQ ID NO: 5.
7. A polypeptide encoded by a human gene that substantially comprises the amino acid sequence as set forth in SEQ ID NO: 2.
- 20 8. A polypeptide encoded by a mouse gene that substantially comprises the amino acid sequence as set forth in SEQ ID NO: 4.
9. A polypeptide encoded by a rat gene that substantially comprises the amino acid sequence as set forth in SEQ ID NO: 6.
- 25 10. An oligonucleotide probe that hybridizes to at least part of the gene according to any one of claims 1 to 6.
11. A recombinant DNA comprising the gene according to any one of claims 1 to 6.
- 30 12. A transformant transformed with the recombinant DNA according to claim 11.
13. A method of producing a polypeptide encoded by a human, mouse and rat gene which method comprises culturing the transformant according to claim 12 and harvesting said polypeptide from the culture.
- 35 14. A monoclonal antibody that specifically reacts



- 43 -

claim 26 said polypeptide having an effect of inhibiting angiogenesis.

ABSTRACT

5 A polypeptide having the amino acid sequence as set forth in SEQ ID NO: 2, 4, or 6, a DNA encoding the same, and an antibody against said polypeptide, and the use thereof. The above amino acid sequence has a homology with chondromodulin-I that has an effect of controlling the growth and differentiation of chondrocytes and inhibiting angiogenesis.

# Fig.1A

hChM-I : human ChM-I

hChM1L : human ChM1L

```
hChM-I      MTENSDKVPIALVGPDDVEFCSPPAYATLTVKPSSPARLLKVGAVVLISGAVLLLFGAIG
hChM1L      -----MAKNPPENCEDCHILNAEAFKSKK--ICKSLKICGLVFGILALTIVLFWG
              *   *   *   *   *   *   *   *   *   *   *   *

hChM-I      AFYFWKGSDSHIYNVHYTMSINGKLQDGSMEIDAGNNLETFKMGSGAEEAIAVNDFQNGI
hChM1L      SKHFWPEVPKKAYDMEHTFYSGEKKKIYMEIDPVTRTEIFRSGNGTDETLEVHDFKNGY
              **   *   *   *   *   *   *   *   *   *   *   *

hChM-I      TGIRFAGGEKCYIKAQVKARIPEVGAVTKQSISSKLEGKIMPVKYEENSLIWVAVDQPVK
hChM1L      TGIYFVGLQKCFIKTQIKV-IPEFSEPEEEID---ENEEITTTTFEQSVIWVPAEKPIE
              *** * * * * * * * * * * * * * * * * * * *

hChM-I      DNSFLS-SKVLELCGDLPIFWLKPTY--KEIQRERREVVVRKIVPTTTKRPHSGPRSNPG
hChM1L      NRDFLKNKILEICDNVTMYWINPTLISVSELQDFEEEGEDLHFPANEKKGIEQNEQWVV
              ** * * * * * * * * * * * * * * * *

hChM-I      AGRLNNETRPSVQEDSQAFNPDNPNYHQEGESMTFDPRLDHEGICCIECRRSYTHCQKIC
hChM1L      PQVKVEKTRHAR----QASEEELPINDYTENGIEFDPMLEDERGYCCIIYCRRGNRYCRRVC
              **   **   *   *   *   *   *   *   *   *   *

hChM-I      EPLGGYYPWPYNYQGCRSACRVIMPCSWWVARILGMV
hChM1L      EPLLGYYPYPYCYQGGRVICRVIMPCNWWVARMLGRV
              *** * * * * * * * * * * * * * * * *
```

## Fig.1B

hChM1L : human ChM1L

mChM1L : mouse ChM1L

rChM1L : rat ChM1L

```
mChM1L      MAKNPPENCEGCHILNAEALKSKKICKSLKICGLVFGILALTLIVLFWGSKHFWPEVSKK
rChM1L      MAKNPPENCEGCHILNAEALKSKKIRKSLKICGLVFGILALTLIVLFWGSKHFWPEVSKK
hChM1L      MAKNPPENCEDCHILNAEAFKSKKICKSLKICGLVFGILALTLIVLFWGSKHFWPEVPPK
*****

mChM1L      TYDMEHTFYSNGEKKKIYMEIDPITRTEIFRSGNGTDETLVHDFKNGYTGIYFVGLQKC
rChM1L      TYGMEHTFYSNGEKKKISMEIDPITRTEIFRSGNGTDETLVHDFKNGYTGIYFVGLQKC
hChM1L      AYDMEHTFYSNGEKKKIYMEIDPVTRTEIFRSGNGTDETLVHDFKNGYTGIYFVGLQKC
* *****

mChM1L      FIKTQIKVIPEFSEPEEEIDENEEITTTFFEQSVIWVPAEKPIENRDFLKNSKILEICDN
rChM1L      FIKTQIKVIPEFSEPEEEIDENEEITTTFFEQSVIWVPAEKPIENRDFLKNSKILEICDN
hChM1L      FIKTQIKVIPEFSEPEEEIDENEEITTTFFEQSVIWVPAEKPIENRDFLKNSKILEICDN
*****

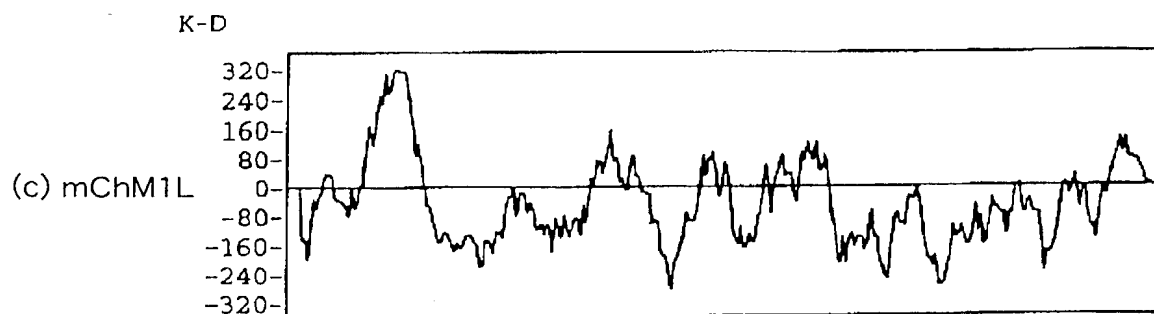
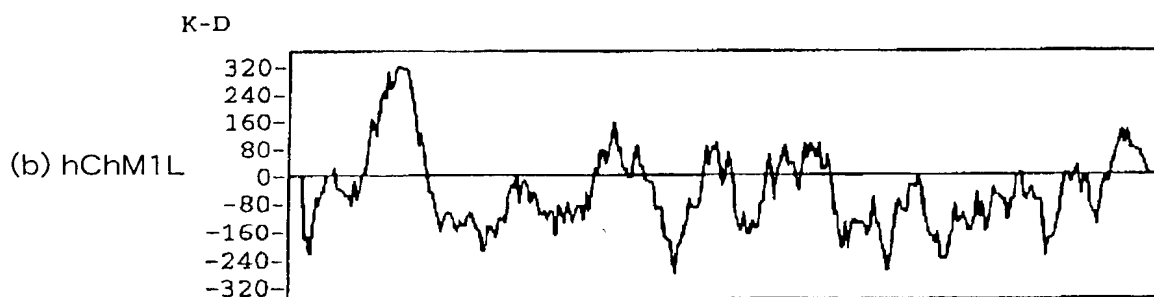
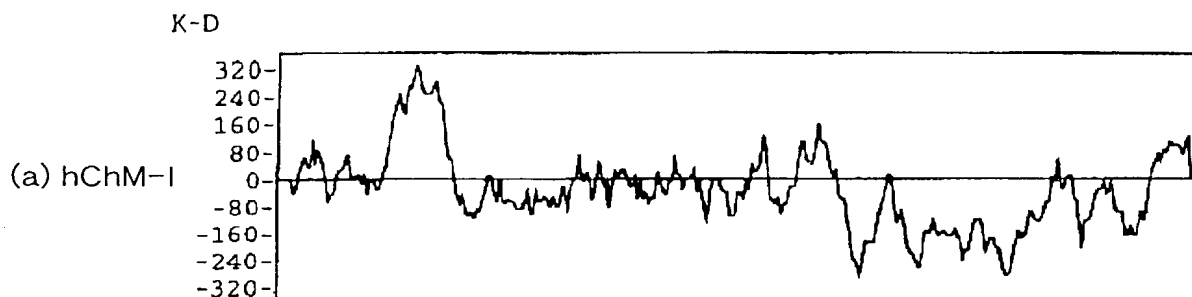
mChM1L      VTMYWINPTLIAVSELQDFEEDGEDLHFPTSEKKGIDQNEQWVVPQVKVEKTRHTRQASE
rChM1L      VTMYWINPTLIAVSELQDFEEDGEDLHFPTSEKKGIDQNEQWVVPQVKVEKTRRTRQASE
hChM1L      VTMYWINPTLISVSELQDFEEEGEDLHFPA NEKKGIEQNEQWVVPQVKVEKTRHARQASE
*****

mChM1L      EDLPINDYTENGIEFDPMLDERGYCCIIYCRGNRYCRRVCEPLLGYYPYPYCYQGGRVIC
rChM1L      EDLPVNDYTENGIEFDPMLDERGYCCIIYCRGNRYCRRVCEPLLGYYPYPYCYQGGRVIC
hChM1L      EELPINDYTENGIEFDPMLDERGYCCIIYCRGNRYCRRVCEPLLGYYPYPYCYQGGRVIC
* * *****

mChM1L      RVIMPCNWWVARMLGRV
rChM1L      RVIMPCNWWVARMLGRV
hChM1L      RVIMPCNWWVARMLGRV
*****
```

## Fig. 2

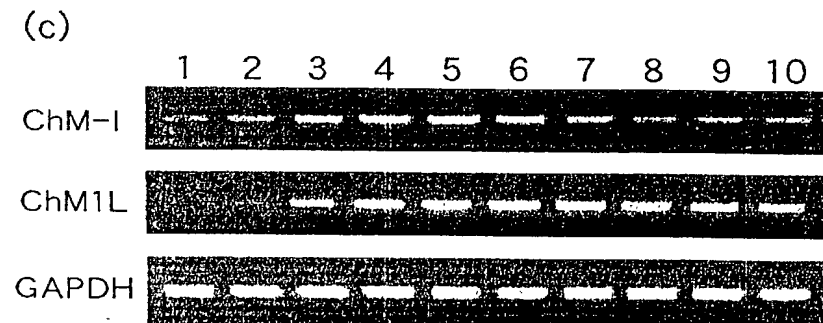
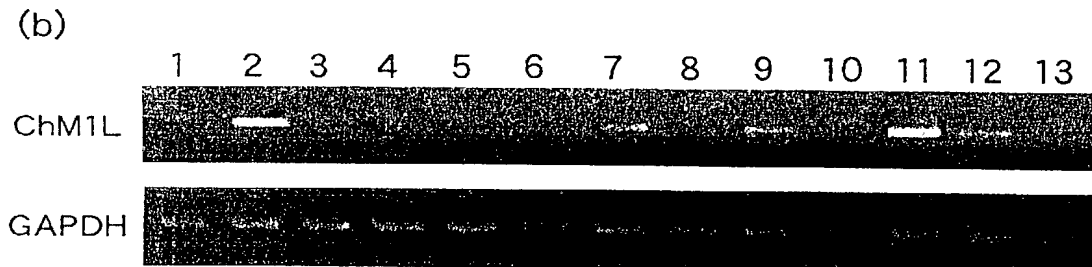
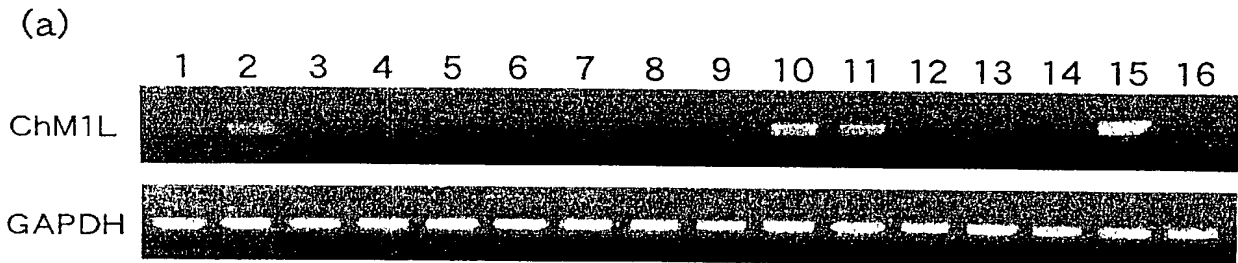
- (a) hChM-I : human ChM-I
- (b) hChM1L : human ChM1L
- (c) mChM1L : mouse ChM1L





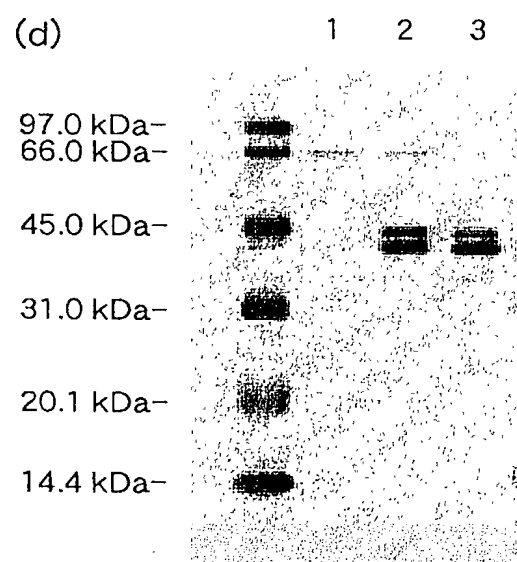
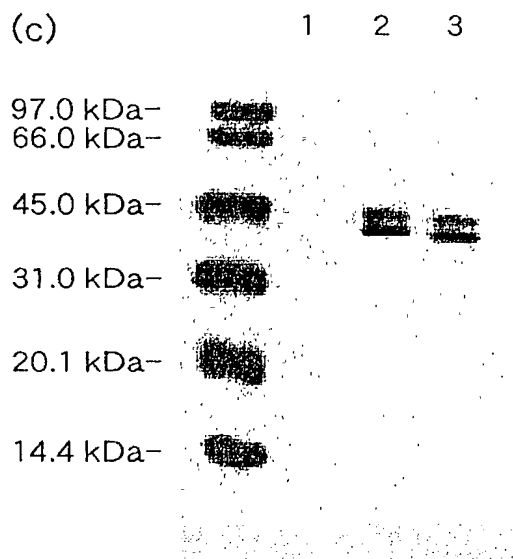
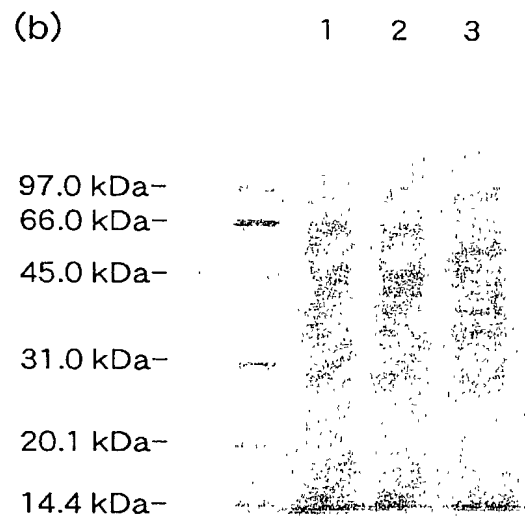
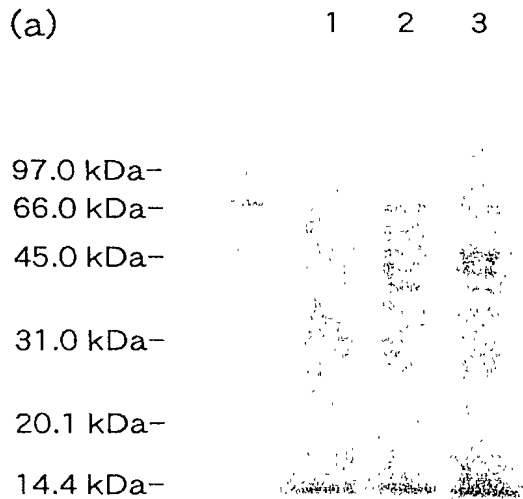
## Fig. 3

- (a) EXPRESSION AT VARIOUS TISSUES OF ADULT (10-WEEK OLD)  
1. BRAIN, 2. EYEBALL, 3. LUNG, 4. THYMUS, 5. HEART, 6. LIVER,  
7. KIDNEY, 8. STOMACH, 9. SPLEEN, 10. SKELETAL MUSCLE, 11.  
WHOLE RIB, 12. FAT, 13. ADRENAL, 14. PITUITARY, 15. THYROID,  
16. INTESTINE
- (b) EXPRESSION AT VARIOUS TISSUES OF FETUS (DAY 17 OF GESTATION)  
1. BRAIN, 2. EYEBALL, 3. LUNG, 4. THYMUS, 5. HEART, 6. LIVER,  
7. KIDNEY, 8. SPLEEN, 9. STOMACH, 10. INTESTINE, 11. WHOLE  
RIB, 12. TRACHEA, 13. PANCREAS
- (c) EXPRESSION DURING THE DEVELOPMENTAL STAGE OF FETUS  
1. DAY 10 OF GESTATION, 2. DAY 11 OF GESTATION, 3. DAY 12 OF  
GESTATION, 4. DAY 13 OF GESTATION, 5. DAY 14 OF GESTATION, 6.  
DAY 15 OF GESTATION, 7. DAY 16 OF GESTATION, 8. DAY 17 OF  
GESTATION, 9. DAY 18 OF GESTATION, 10. DAY OF CHILDBIRTH



# Fig. 4

- (a) SDS-PAGE: 1. MOCK, 2. HUMAN ChM1L, 3. MOUSE ChM1L
- (b) SDS-PAGE: 1. MOCK, 2. HUMAN ChM1L(His), 3. MOUSE ChM1L(His)
- (c) WESTERN BLOT (DETECTION WITH ANTI-PEPTIDE ANTIBODY):  
 1. MOCK, 2. HUMAN ChM1L, 3. MOUSE ChM1L
- (d) WESTERN BLOT (DETECTION WITH ANTI-His TAG ANTIBODY):  
 1. MOCK, 2. HUMAN ChM1L(His), 3. MOUSE ChM1L(His)



## Fig. 5

1. Mock
2. soluble human ChM1L

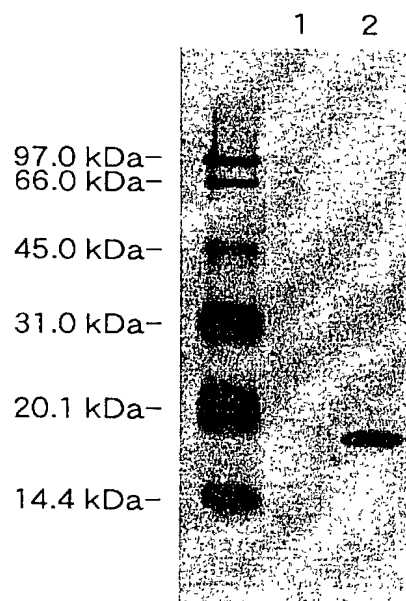
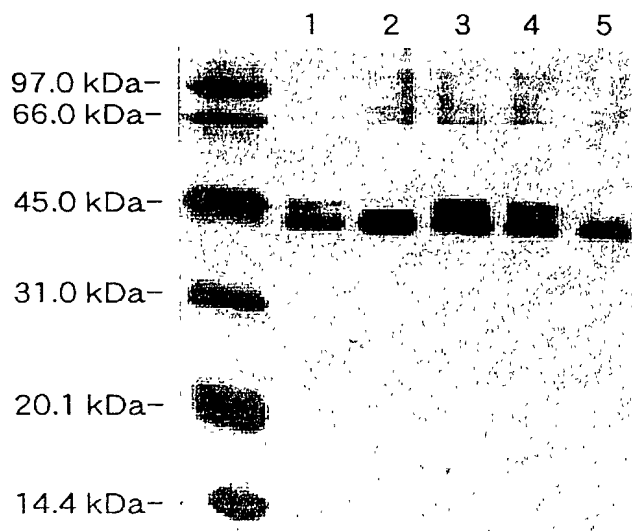


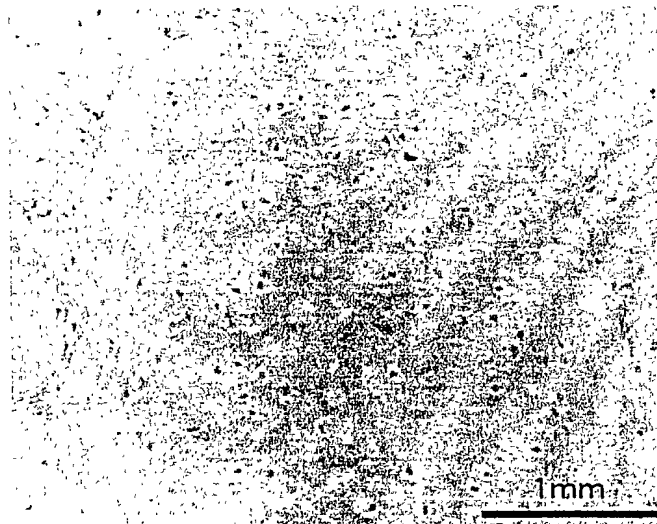
Fig. 6

1. NON-TREATED, 2. TREATED WITH NANase II + O-GLYCOSIDASE DS + PNGase F, 3. TREATED WITH NANase II, 4. TREATED WITH O-GLYCOSIDASE DS, AND 5. TREATED WITH PNGase F



## Fig.7

(a) RABBIT IgG



(b) ANTI-ChM1L PEPTIDE ANTIBODY

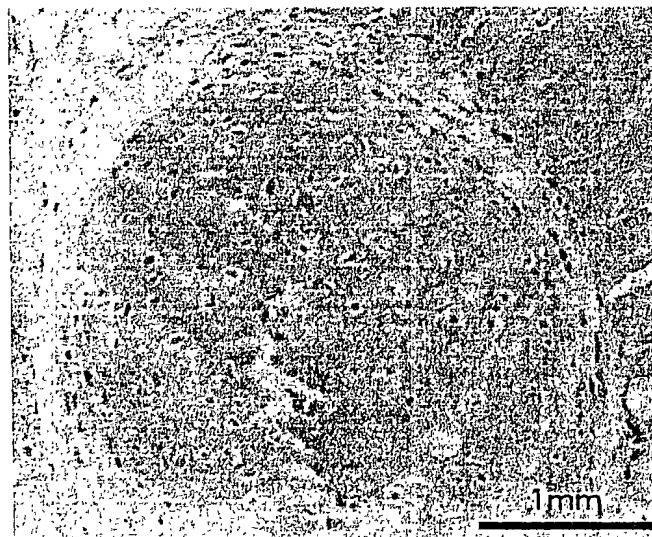


Fig. 8

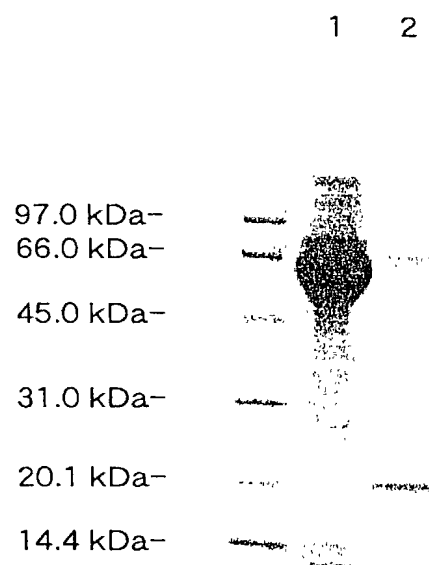


Fig. 9

# Declaration and Power of Attorney for Patent Application

## 特許出願宣言書および委任状

### Japanese Language Declaration

#### 日本語宣言書

私は下記発明者として以下の通り宣言します：

私の住所、郵送先、および国籍は私の氏名の後に記載された通りです。

下記名称の発明に関し請求範囲に記載され特許出願がされている発明内容につき、私が最初、最先かつ唯一の発明者（下記氏名が一つのみの場合）であるか、あるいは最初、最先かつ共同発明者（下記氏名が複数の場合）であると信じます。

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\_\_\_\_\_  
\_\_\_\_\_

下記項目に×印が付いている場合を除き、上記発明の明細書は本書に添付されます。

☐ 上記発明は米国出願番号あるいはPCT国際出願番号（確認番号\_\_\_\_\_）として\_\_\_\_年\_\_月\_\_日に出願され、\_\_\_\_年\_\_月\_\_日に補正されました（該当する場合）。

私は特許請求範囲を含み上述の補正で補正された前記明細書の内容を検討し、理解していることをここに表明します。

私は連邦規則法典第 37 編 1 条 56 項に定義される特許性に肝要な情報について開示義務があることを認めます。

As a below named inventor, I hereby declare that:

My residence, mailing address and citizenship are as stated next to my name.

I believe I am the original, first and sole inventor (if only one name is listed below) or an original, first and joint inventor (if plural names are listed below) of the subject matter which is claimed and for which a patent is sought on the invention entitled

A Novel polypeptide and gene encoding  
the same

\_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_

the specification of which is attached hereto unless the following box is checked:

☒ was filed on September 29, 2000  
as United States Application Number or  
PCT International Application Number  
PCT/JP00/06804 (Conf. No. \_\_\_\_\_)  
and was amended on \_\_\_\_\_  
(if applicable).

I hereby state that I have reviewed and understand the contents of the above identified specification, including the claims, as amended by any amendment referred to above.

I acknowledge the duty to disclose information which is material to patentability as defined in Title 37, Code of Federal Regulations, § 1.56.



# Japanese Language Declaration

## 日本語宣言書

私は米国法典第 35 編 119 条(a)-(d)あるいは 365 条(b)に基づき特許あるいは発明者証書の下記外国出願、または 365 条(a)に基づき米国以外の少なくとも 1 ケ国を指定した下記 PCT 外国出願についての外国優先権をここに主張するとともに、下記項目に×印を付けることにより優先権を主張する出願以前の出願日を有する特許あるいは発明者証書の外国出願あるいは PCT 外国出願を示します。

I hereby claim foreign priority under Title 35, United States Code, § 119(a)-(d) or § 365(b) of any foreign application(s) for patent or inventor's certificate, or § 365(a) of any PCT International application which designated at least one country other than the United States, listed below, and have also identified below, by checking the box, any foreign application for patent or inventor's certificate, or PCT International application having a filing date before that of the application on which priority is claimed.

Prior foreign application(s)

外国での先行出願

Priority Claimed

優先権の主張

Yes No  
有り無し

11-275947(Pat.Appln.) Japan  
(Number) (Country)  
(番号) (国名)

29 / September / 1999

(Day/Month/Year Filed)  
(出願年月日)

☒ ☐

(Number) (Country)  
(番号) (国名)

(Day/Month/Year Filed)  
(出願年月日)

私は米国法典第 35 編 119 条(e)に基づき下記の米国仮特許の利益をここに主張します。

I hereby claim the benefit under Title 35, United States Code, § 119(e) of any United States provisional application(s) listed below.

(Application No.)  
(出願番号)

(Filing Date)  
(出願日)

(Application No.)  
(出願番号)

(Filing Date)  
(出願日)

私は米国法典第 35 編 120 条に基づき下記米国特許出願、あるいは 365 条(c)に基づき米国を指定する下記 PCT 国際特許出願の利益をここに主張し、本特許出願内特許請求範囲の各項目の内容が米国法典第 35 編 112 条の最初の項に規定される方法により先行米国あるいは PCT 国際特許出願で開示されていない限りにおいて連邦規則法典第 37 編 1 条 56 項に定義される特許性に肝要で、先行特許出願の出願日から本特許出願の国内あるいは PCT の出願日までの間に入手された情報について開示義務があることを認めます。

I hereby claim the benefit under Title 35, United States Code, § 120 of any United States application(s), or § 365(c) of any PCT International application designating the United States, listed below and, insofar as the subject matter of each of the claims of this application is not disclosed in the prior United States or PCT International application in the manner provided by the first paragraph of Title 35, United States Code, § 112, I acknowledge the duty to disclose information which is material to patentability as defined in Title 37, Code of Federal Regulations, § 1.56 which became available between the filing date of the prior application and the national or PCT International filing date of this application.

(Application No.)  
(出願番号)

(Filing Date)  
(出願日)

(Status: patented, pending, abandoned)  
(状態: 特許成立済、係属中、放棄済)

(Application No.)  
(出願番号)

(Filing Date)  
(出願日)

(Status: patented, pending, abandoned)  
(状態: 特許成立済、係属中、放棄済)

私は本宣言書内で私自身の知識に基づいてなされたすべての陳述が真実であり、情報および信ずるところに基づいてなされたすべての陳述が真実であると信じられていることをここに宣言し、さらに故意になされた虚偽の陳述等々は米国法典第 18 編 1001 条に基づき罰金あるいは拘禁または両方による処罰にあたり、またかような故意による虚偽の陳述はそれに基づく特許出願あるいは成立特許の有効性を危うくする可能性があることを認識した上でこれらの陳述をなしたことを宣言します。

I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

# Japanese Language Declaration

## 日本語宣言書

委任状：私は下記の米国特許商標局（USPTO）顧客番号のもとに記載される SUGHRUE MION 法律事務所のすべての弁護士を、同顧客番号のもとに記載される個々の弁護士は SUGHRUE MION 法律事務所のための自由裁量に基づき変更され得ることを認識した上で、本特許出願の手続きおよびそれに関わる特許商標局との業務を遂行する弁護士として指名し、本特許出願に関するすべての通信が同 USPTO 顧客番号のもとに提出された住所宛に送付されることを要請します。

POWER OF ATTORNEY: I hereby appoint all attorneys of SUGHRUE MION, PLLC who are listed under the USPTO Customer Number shown below as my attorneys to prosecute this application and to transact all business in the Patent and Trademark Office connected therewith, recognizing that the specific attorneys listed under that Customer Number may be changed from time to time at the sole discretion of SUGHRUE MION, PLLC, and request that all correspondence about the application be addressed to the address filed under the same USPTO Customer Number.



23373

PATENT TRADEMARK OFFICE

電話連絡は下記へ：(名前および電話番号)

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SUGHRUE MION, PLLC  
(202) 293-7060

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Full name of sole or first inventor 唯一あるいは第一の発明者名 <u>Kei Yamana</u>	
Inventor's signature 発明者の署名 <u>Kei Yamana</u>	Date 日付 March 12, 2002
Residence 住所 Hino-shi, Tokyo, Japan JPX	
Citizenship 国籍 Japanese	
Mailing Address 郵送先 c/o TEIJIN LIMITED, Tokyo Research Center, 3-2, Asahigaoka 4-chome, Hino-shi, Tokyo 191-0065, Japan	
Full name of second joint inventor, if any 第二の共同発明者名(該当する場合) <u>Yukimi Nagasawa</u>	
Second inventor's signature 第二発明者の署名 <u>Yukimi Nagasawa</u>	Date 日付 March 12, 2002
Residence 住所 Hino-shi, Tokyo, Japan JPX	
Citizenship 国籍 Japanese	
Mailing Address 郵送先 c/o TEIJIN LIMITED, Tokyo Research Center, 3-2, Asahigaoka 4-chome, Hino-shi, Tokyo 191-0065, Japan	



## SEQUENCE LISTING

&lt;110&gt; Teijin Limited

&lt;120&gt; A NOVEL POLYPEPTIDE AND GENE ENCODING THE SAME

&lt;130&gt; Q69170

&lt;150&gt; PCT/JP00/06804

&lt;151&gt; 2000-09-29

&lt;150&gt; JP 11-275947

&lt;151&gt; 1999-09-29

&lt;160&gt; 25

&lt;170&gt; PatentIn version 3.1

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 Met Ser Ala Leu Leu Ile Leu Ala Leu Val Gly Ala Ala Val Ala Asp  
 1 5 10 15  
 tac aaa gac gat gac gac aag ctg gaa ttc gat gag gga gaa gat ctt 96  
 Tyr Lys Asp Asp Asp Lys Leu Glu Phe Asp Glu Gly Glu Asp Leu  
 20 25 30  
 cac ttt cct gcc aac gaa aaa aaa ggg att gaa caa aat gaa cag tgg 144  
 His Phe Pro Ala Asn Glu Lys Lys Gly Ile Glu Gln Asn Glu Gln Trp  
 35 40 45  
 gtg gtc cct caa gtg aaa gta gag aag acc cgt cac gcc aga caa gca 192  
 Val Val Pro Gln Val Lys Val Glu Lys Thr Arg His Ala Arg Gln Ala  
 50 55 60  
 agt gag gaa gaa ctt cca ata aat gac tat act gaa aat gga ata gaa 240  
 Ser Glu Glu Glu Leu Pro Ile Asn Asp Tyr Thr Glu Asn Gly Ile Glu  
 65 70 75 80  
 ttt gat ccc atg ctg gat gag aga ggt tat tgt tgt att tac tgc cgt 288  
 Phe Asp Pro Met Leu Asp Glu Arg Gly Tyr Cys Cys Ile Tyr Cys Arg  
 85 90 95  
 cga ggc aac cgc tat tgc cgc cgc gtc tgt gaa cct tta cta ggc tac 336  
 Arg Gly Asn Arg Tyr Cys Arg Arg Val Cys Glu Pro Leu Leu Gly Tyr  
 100 105 110  
 tac cca tat cca tac tgc tac caa gga gga cga gtc atc tgt cgt gtc 384  
 Tyr Pro Tyr Pro Tyr Cys Tyr Gln Gly Gly Arg Val Ile Cys Arg Val  
 115 120 125  
 atc atg cct tgt aac tgg tgg gtg gcc cgc atg ctg ggg agg gtc taa 432  
 Ile Met Pro Cys Asn Trp Trp Val Ala Arg Met Leu Gly Arg Val  
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<223> signal sequence of preprotrypsin, FLAG peptide and C terminal region of ChM1L

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Met Ser Ala Leu Leu Ile Leu Ala Leu Val Gly Ala Ala Val Ala Asp  
1 5 10 15

Tyr Lys Asp Asp Asp Asp Lys Leu Glu Phe Asp Glu Gly Glu Asp Leu  
20 25 30

His Phe Pro Ala Asn Glu Lys Lys Gly Ile Glu Gln Asn Glu Gln Trp  
35 40 45

Val Val Pro Gln Val Lys Val Glu Lys Thr Arg His Ala Arg Gln Ala  
50 55 60

Ser Glu Glu Glu Leu Pro Ile Asn Asp Tyr Thr Glu Asn Gly Ile Glu  
65 70 75 80

Phe Asp Pro Met Leu Asp Glu Arg Gly Tyr Cys Cys Ile Tyr Cys Arg  
85 90 95

Arg Gly Asn Arg Tyr Cys Arg Arg Val Cys Glu Pro Leu Leu Gly Tyr  
100 105 110

Tyr Pro Tyr Pro Tyr Cys Tyr Gln Gly Gly Arg Val Ile Cys Arg Val  
115 120 125

Ile Met Pro Cys Asn Trp Trp Val Ala Arg Met Leu Gly Arg Val  
130 135 140

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<212> DNA

<213> Mus musculus

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tcagccatga cagagaactc a

21

<210> 25

<211> 21

<212> DNA

<213> Mus musculus

<400> 25

ttacaccatg cccaagatgc g

21